

SEQUENCE LISTING

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<120> Fv Antibody Construct Comprising Binding Sites For a CD16 Receptor and a CD30 Surface Protein

<130> 4121-135

<140> US 10/049,404

<141> 2002-02-05

<150> PCT/DE00/02589

<151> 2000-08-02

<150> DE 199 37 264

<151> 1999-08-06

<160> 11

<170> PatentIn version 3.1

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<211> 4570

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Trp Ile Gly Asp Ile Tyr Pro Gly Gly Gly Tyr Thr Asn Tyr Asn Glu
    50          55          60

Lys Phe Lys Gly Lys Ala Thr Val Thr Ala Asp Thr Ser Ser Arg Thr
    65          70          75          80

Ala Tyr Val Gln Val Arg Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr
    85          90          95

Phe Cys Ala Arg Ser Ala Ser Trp Tyr Phe Asp Val Trp Gly Ala Arg
    100         105         110

Thr Thr Val Thr Val Ser Ser Ala Lys Thr Thr Pro Lys Leu Gly Gly
    115         120         125

Asp Ile Glu Leu Thr Gln Ser Pro Lys Phe Met Ser Thr Ser Val Gly
    130         135         140

Asp Arg Val Asn Val Thr Tyr Lys Ala Ser Gln Asn Val Gly Thr Asn
    145         150         155         160

Val Ala Trp Phe Gln Gln Lys Pro Gly Gln Ser Pro Lys Val Leu Ile
    165         170         175

Tyr Ser Ala Ser Tyr Arg Tyr Ser Gly Val Pro Asp Arg Phe Thr Gly
    180         185         190

Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Asn Val Gln Ser
    195         200         205

Glu Asp Leu Ala Glu Tyr Phe Cys Gln Gln Tyr His Thr Tyr Pro Leu
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Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg Ala Asp Ala Ala
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 20 25 30

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 35 40 45

Trp Ile Gly Tyr Ile Asn Pro Ser Ser Gly Tyr Ser Asp Tyr Asn Gln
 50 55 60

Asn Phe Lys Gly Lys Thr Thr Leu Thr Ala Asp Lys Ser Ser Asn Thr
 65 70 75 80

Ala Tyr Met Gln Leu Asn Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr
 85 90 95

Tyr Cys Ala Arg Arg Ala Asp Tyr Gly Asn Tyr Glu Tyr Thr Trp Phe
 100 105 110

Ala Tyr Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Ala Lys Thr
 115 120 125

Thr Pro Lys Leu Gly Gly Asp Ile Gln Ala Val Val Thr Gln Glu Ser
 130 135 140

Ala Leu Thr Thr Ser Pro Gly Glu Thr Val Thr Leu Thr Cys Arg Ser
 145 150 155 160

Asn Thr Gly Thr Val Thr Thr Ser Asn Tyr Ala Asn Trp Val Gln Glu
 165 170 175

Lys Pro Asp His Leu Phe Thr Gly Leu Ile Gly His Thr Asn Asn Arg
 180 185 190

Ala Pro Gly Val Pro Ala Arg Phe Ser Gly Ser Leu Ile Gly Asp Lys
 195 200 205

Ala Ala Leu Thr Ile Thr Gly Ala Gln Thr Glu Asp Glu Ala Ile Tyr
 210 215 220

Phe Cys Ala Leu Trp Tyr Asn Asn His Trp Val Phe Gly Gly Gly Thr
 225 230 235 240

Lys Leu Thr Val Leu Gly Gln Pro Lys Ser Ala Ala Ala Gly Ser Glu
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Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn Ser His His His His His
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 <400> 5
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 <210> 7
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